	(2)	INFORMATION FOR SEQ ID NO:39:			
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 21 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:39:		
	AAC	AGCTA:	IG ACCATGATTA C	21	
	(2)	2) INFORMATION FOR SEQ ID NO:40:			
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 60 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:40:		
0	GTT	CTCTG	CT CTCTGGTCGC TGTCTCGCTT GTGAAACAAG CGAGACAGCG TGGTCTCTCG	60	
	(2)	INFO	RMATION FOR SEQ ID NO:41:		
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 15 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:41:		
	CGAGAGACCA CGCTG				
	(2)	INFO	RMATION FOR SEQ ID NO:42:		
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 52 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:42:		
	CCT	TTCGC	TT TCTTCCCTTC CTTTCTCGCC ACGTTCGCCG GCTTTCCCCG TC	52	

	(2)	INFO	RMATION FOR SEQ ID NO:43:		
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 26 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:43:		
	AGA	AAGGA	AG GGAAGAAGC GAAAGG	26	
	(2)	INFO	RMATION FOR SEQ ID NO:44:		
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 21 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:44:		
	GAC	GGGGA	AA GCCGGCGAAC G	21	
	(2)	INFO	RMATION FOR SEQ ID NO:45:		
7		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:45:		
	GAAAGCCGGC GAACGTGGCG				
	(2)	INFO	RMATION FOR SEQ ID NO:46:		
		(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 21 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
		(ii)	MOLECULE TYPE: DNA (genomic)		
		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:46:		
	GGC	GAAC	GTG GCGAGAAAGG A	21	

(2)	INFORMATION FOR SEQ ID NO:47:			
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 42 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
	(ii)	MOLECULE TYPE: DNA (genomic)		
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:47:		
CCT	TTCGC'	TT TCTTCCCTTC CTTTCTCGCC ACGTTCGCCG GC	42	
(2)	INFO	RMATION FOR SEQ ID NO:48:		
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 42 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
	(ii)	MOLECULE TYPE: DNA (genomic)		
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:48:		
CCT	TTCGC	TC TCTTCCCTTC CTTTCTCGCC ACGTTCGCCG GC	42	
(2)	INFO	RMATION FOR SEQ ID NO:49:		
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 27 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
	(ii)	MOLECULE TYPE: DNA (genomic)		
,	mod h	FEATURE:  (A) NAME/KEY: modified_base  (B) LOCATION: 8  (C) IDENTIFICATION METHOD: experimental  (D) OTHER INFORMATION: /evidence= EXPERIMENTAL  case= OTHER  "The A residue at this position is 2'-O-methyladenosine."		
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:49:		
AGA	AAAGG2	AAG GGAAGAAAGC GAAAGGT	27	
(2)	INF	ORMATION FOR SEQ ID NO:50:		
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
	(11	) MOLECULE TYPE: DNA (genomic)		